

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/664,506A

Source: IFW/6

Date Processed by STIC: 8/9/05

ENTERED

BEST AVAILABLE COPY

**CRF Errors Edited by the STIC Systems
Branch**

Serial Number: 10/664,506A

CRF Edit Date: 8/10/05
Edited by: W

___ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

___ Corrected the SEQ ID NO. Sequence numbers edited were:

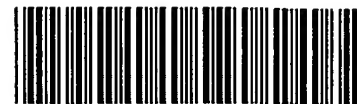
___ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

___ Deleted: / invalid beginning/end-of-file text ; ___ page numbers

___ Inserted mandatory headings/numeric identifiers, specifically:

___ Moved responses to same line as heading/numeric identifier, specifically:

___ Other:



IFW16

RAW SEQUENCE LISTING

DATE: 08/10/2005

PATENT APPLICATION: US/10/664,506A

TIME: 13:55:23

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08102005\J664506A.raw

4 <110> APPLICANT: Meyers, Rachel
 5 Cook, William James
 6 Williamson, Mark
 7 Rudolph-Owen, Laura A.
 8 Gimeno, Ruth
 10 <120> TITLE OF INVENTION: 21481, A NOVEL DEHYDROGENASE MOLECULE
 11 AND USES THEREFOR
 14 <130> FILE REFERENCE: MPI00-079P1RCP2CN1M
 16 <140> CURRENT APPLICATION NUMBER: 10/664,506A
 17 <141> CURRENT FILING DATE: 2003-09-17
 19 <150> PRIOR APPLICATION NUMBER: 09/838,561
 20 <151> PRIOR FILING DATE: 2001-04-18
 22 <150> PRIOR APPLICATION NUMBER: 09/816,760
 23 <151> PRIOR FILING DATE: 2001-03-23
 25 <150> PRIOR APPLICATION NUMBER: 09/634,955
 26 <151> PRIOR FILING DATE: 2000-08-08
 28 <150> PRIOR APPLICATION NUMBER: 60/192,002
 29 <151> PRIOR FILING DATE: 2000-03-24
 31 <160> NUMBER OF SEQ ID NOS: 34
 33 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 35 <210> SEQ ID NO: 1
 36 <211> LENGTH: 2660
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Homo sapiens
 40 <220> FEATURE:
 41 <221> NAME/KEY: CDS
 42 <222> LOCATION: (63)...(2468)
 44 <221> NAME/KEY: misc_feature
 45 <222> LOCATION: 6, 8
 46 <223> OTHER INFORMATION: n = A,T,C or G
 W--> 48 <400> 1
 W--> 49 cctttntnrc cacgcgtccg agagcgcgcc gacgtcttcg cggaaagcgt tcggggtagg 60
 50 cg atg gct gcg acg cgt gca ggg ccc cgc gcc cgc gag atc ttc acc 107
 51 Met Ala Ala Thr Arg Ala Gly Pro Arg Ala Arg Glu Ile Phe Thr
 52 1 5 10 15
 54 tcg ctg gag tac gga ccg gtg ccg gag agc cac gca tgc gca ctg gcc 155
 55 Ser Leu Glu Tyr Gly Pro Val Pro Glu Ser His Ala Cys Ala Leu Ala
 56 20 25 30
 58 tgg ctg gac acc cag gac cgg tgc ttg ggc cac tat gtg aat ggg aag 203
 59 Trp Leu Asp Thr Gln Asp Arg Cys Leu Gly His Tyr Val Asn Gly Lys
 60 35 40 45
 62 tgg tta aag cct gaa cac aga aat tca gtg cct tgc cag gat ccc atc 251
 63 Trp Leu Lys Pro Glu His Arg Asn Ser Val Pro Cys Gln Asp Pro Ile

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DATE: 08/10/2005

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Input Set : A:\PTO.AMC.txt

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| | | | | |
|-----|---|------|-----|-----|
| 64 | 50 | 55 | 60 | |
| 66 | aca gga gag aac ttg gcc agt tgc ctg cag gca cag gcc gag gat gtg | 299 | | |
| 67 | Thr Gly Glu Asn Leu Ala Ser Cys Leu Gln Ala Gln Ala Glu Asp Val | | | |
| 68 | 65 | 70 | 75 | |
| 70 | gct gca gcc gtg gag gca gcc agg atg gca ttt aag ggc tgg agt gcg | 347 | | |
| 71 | Ala Ala Ala Val Glu Ala Ala Arg Met Ala Phe Lys Gly Trp Ser Ala | | | |
| 72 | 80 | 85 | 90 | 95 |
| 74 | cac ccc ggc gtc gtc cgg gcc cag cac ctg acc agg ctg gcc gag gtg | 395 | | |
| 75 | His Pro Gly Val Val Arg Ala Gln His Leu Thr Arg Leu Ala Glu Val | | | |
| 76 | 100 | 105 | 110 | |
| 78 | atc cag aag cac cag cgg ctg ctg tgg acc ctg gaa tcc ctg gtg act | 443 | | |
| 79 | Ile Gln Lys His Gln Arg Leu Leu Trp Thr Leu Glu Ser Leu Val Thr | | | |
| 80 | 115 | 120 | 125 | |
| 82 | ggg cgg gct gtt cga gag gtt cga gac ggg gac gtc cag ctg gcc cag | 491 | | |
| 83 | Gly Arg Ala Val Arg Glu Val Arg Asp Gly Asp Val Gln Leu Ala Gln | | | |
| 84 | 130 | 135 | 140 | |
| 86 | cag ctg ctc cac tac cat gca atc cag gca tcc acc cag gag gag gca | 539 | | |
| 87 | Gln Leu Leu His Tyr His Ala Ile Gln Ala Ser Thr Gln Glu Glu Ala | | | |
| 88 | 145 | 150 | 155 | |
| 90 | ctg gca ggc tgg gag ccc atg gga gta att ggc ctc atc ctg cca ccc | 587 | | |
| 91 | Leu Ala Gly Trp Glu Pro Met Gly Val Ile Gly Leu Ile Leu Pro Pro | | | |
| 92 | 160 | 165 | 170 | 175 |
| 94 | aca ttc tcc ttc ctt gag atg atg tgg agg att tgc cct gcc ctg gct | 635 | | |
| 95 | Thr Phe Ser Phe Leu Glu Met Met Trp Arg Ile Cys Pro Ala Leu Ala | | | |
| 96 | 180 | 185 | 190 | |
| 98 | gtg ggc tgc acc gtg gtg gcc ctc gtg ccc ccg gcc tcc ccg gcg ccc | 683 | | |
| 99 | Val Gly Cys Thr Val Val Ala Leu Val Pro Pro Ala Ser Pro Ala Pro | | | |
| 100 | 195 | 200 | 205 | |
| 102 | ctc ctc ctg gcc cag ctg gcg ggg gag ctg ggc ccc ttc ccg gga atc | 731 | | |
| 103 | Leu Leu Leu Ala Gln Leu Ala Gly Glu Leu Gly Pro Phe Pro Gly Ile | | | |
| 104 | 210 | 215 | 220 | |
| 106 | ctg aat gtc gtc agt ggc cct gcg tcc ctg gtg ccc atc ctg gcc tcc | 779 | | |
| 107 | Leu Asn Val Val Ser Gly Pro Ala Ser Leu Val Pro Ile Leu Ala Ser | | | |
| 108 | 225 | 230 | 235 | |
| 110 | cag cct gga atc cgg aag gtg gcc ttc tgc gga gcc ccg gag gaa ggg | 827 | | |
| 111 | Gln Pro Gly Ile Arg Lys Val Ala Phe Cys Gly Ala Pro Glu Glu Gly | | | |
| 112 | 240 | 245 | 250 | 255 |
| 114 | cgt gcc ctt cga cgg agc ctg gcg gga gag tgt gcg gag ctg ggc ctg | 875 | | |
| 115 | Arg Ala Leu Arg Arg Ser Leu Ala Gly Glu Cys Ala Glu Leu Gly Leu | | | |
| 116 | 260 | 265 | 270 | |
| 118 | gcg ctg ggg acg gag tgc ctg ctg ctg ctg acg gac acg gcg gac gta | 923 | | |
| 119 | Ala Leu Gly Thr Glu Ser Leu Leu Leu Leu Thr Asp Thr Ala Asp Val | | | |
| 120 | 275 | 280 | 285 | |
| 122 | gac tgc gcc gtg gag ggt gtc gtg gac gcc gcc tgg tcc gac cgc ggc | 971 | | |
| 123 | Asp Ser Ala Val Glu Gly Val Val Asp Ala Ala Trp Ser Asp Arg Gly | | | |
| 124 | 290 | 295 | 300 | |
| 126 | ccg ggt ggc ctc agg ctc ctc atc cag gag tct gtg tgg gat gaa gcc | 1019 | | |
| 127 | Pro Gly Gly Leu Arg Leu Leu Ile Gln Glu Ser Val Trp Asp Glu Ala | | | |
| 128 | 305 | 310 | 315 | |

RAW SEQUENCE LISTING

DATE: 08/10/2005

PATENT APPLICATION: US/10/664,506A

TIME: 13:55:23

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08102005\J664506A.raw

```

130 atg aga cgg ctg cag gag cgg atg ggg cgg ctt cgg agt ggc cga ggg 1067
131 Met Arg Arg Leu Gln Glu Arg Met Gly Arg Leu Arg Ser Gly Arg Gly
132 320 325 330 335
134 ctg gat ggg gcc gtg gac atg ggg gcc cgg ggg gct gcc gca tgt gac 1115
135 Leu Asp Gly Ala Val Asp Met Gly Ala Arg Gly Ala Ala Ala Cys Asp
136 340 345 350
138 ctg gtc cag cgc ttt gtg cgt gag gcc cag agc cag ggt gca cag gtg 1163
139 Leu Val Gln Arg Phe Val Arg Glu Ala Gln Ser Gln Gly Ala Gln Val
140 355 360 365
142 ttc cag gct ggt gat gtg cct tcg gaa cgc cca ttc tat ccc cca acc 1211
143 Phe Gln Ala Gly Asp Val Pro Ser Glu Arg Pro Phe Tyr Pro Pro Thr
144 370 375 380
146 ttg gtc tcc aac ctg ccc cca gcc tcc cca tgt gcc cag gtg gag gtg 1259
147 Leu Val Ser Asn Leu Pro Pro Ala Ser Pro Cys Ala Gln Val Glu Val
148 385 390 395
150 ccg tgg cct gtg gtc gtg gcc tcc ccc ttc cgc aca gcc aag gag gca 1307
151 Pro Trp Pro Val Val Val Ala Ser Pro Phe Arg Thr Ala Lys Glu Ala
152 400 405 410 415
154 ctg ttg gtg gcc aac ggg acg ccc cgc ggg ggc agc gcc agt gtg tgg 1355
155 Leu Leu Val Ala Asn Gly Thr Pro Arg Gly Gly Ser Ala Ser Val Trp
156 420 425 430
158 agc gag agg ctg ggg cag gcg ctg gag ctg ggc tat ggg ctc cag gtg 1403
159 Ser Glu Arg Leu Gly Gln Ala Leu Glu Leu Gly Tyr Gly Leu Gln Val
160 435 440 445
162 ggc act gtc tgg atc aac gcc cac ggc ctc aga gac cct tcg gtg ccc 1451
163 Gly Thr Val Trp Ile Asn Ala His Gly Leu Arg Asp Pro Ser Val Pro
164 450 455 460
166 aca ggc ggc tgc aag gag agt ggg tgt tcc tgg cac ggg ggc cca gac 1499
167 Thr Gly Gly Cys Lys Glu Ser Gly Cys Ser Trp His Gly Gly Pro Asp
168 465 470 475
170 ggg ctg tat gag tat ctg cgg ccc tca ggg acc cct gcc cgg ctg tcc 1547
171 Gly Leu Tyr Glu Tyr Leu Arg Pro Ser Gly Thr Pro Ala Arg Leu Ser
172 480 485 490 495
174 tgc ctc tcc aag aac ctg aac tat gac acc ttt ggc ctc gct gtg ccc 1595
175 Cys Leu Ser Lys Asn Leu Asn Tyr Asp Thr Phe Gly Leu Ala Val Pro
176 500 505 510
178 tca acc ctg ccg gct ggg cct gaa ata ggg ccc agc cca gca ccc ccc 1643
179 Ser Thr Leu Pro Ala Gly Pro Glu Ile Gly Pro Ser Pro Ala Pro Pro
180 515 520 525
182 tat ggg ctc ttc gtt ggg ggc cgt ttc cag gct cct ggg gcc cga agc 1691
183 Tyr Gly Leu Phe Val Gly Gly Arg Phe Gln Ala Pro Gly Ala Arg Ser
184 530 535 540
186 tcc agg ccc atc cgg gat tcg tct ggc aat ctc cat ggc tac gtg gct 1739
187 Ser Arg Pro Ile Arg Asp Ser Ser Gly Asn Leu His Gly Tyr Val Ala
188 545 550 555
190 gag ggt gga gcc aag gac atc cga ggt gct gtg gag gcc gct cac cag 1787
191 Glu Gly Gly Ala Lys Asp Ile Arg Gly Ala Val Glu Ala Ala His Gln
192 560 565 570 575
194 gct ttc cct ggc tgg gcg ggc cag tcc cca gga gcc cgg gca gcc ctg 1835

```

RAW SEQUENCE LISTING

DATE: 08/10/2005

PATENT APPLICATION: US/10/664,506A

TIME: 13:55:23

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08102005\J664506A.raw

```

195 Ala Phe Pro Gly Trp Ala Gly Gln Ser Pro Gly Ala Arg Ala Ala Leu
196      580      585      590
198 ctg tgg gcc ctg gcg gct gca ctg gag cgc cgg aag tct acc ctg gcc 1883
199 Leu Trp Ala Leu Ala Ala Ala Leu Glu Arg Arg Lys Ser Thr Leu Ala
200      595      600      605
202 tca agg ctg gag agg cag gga gcg gag ctg aag gct gcg gag gcg gag 1931
203 Ser Arg Leu Glu Arg Gln Gly Ala Glu Leu Lys Ala Ala Glu Ala Glu
204      610      615      620
206 gtg gag ctg agc gca aga cga ctt cgg gcg tgg ggg gcc cgg gtg cag 1979
207 Val Glu Leu Ser Ala Arg Arg Leu Arg Ala Trp Gly Ala Arg Val Gln
208      625      630      635
210 gcc caa ggc cac acc ctg cag gta gcc ggg ctg aga ggc cct gtg ctg 2027
211 Ala Gln Gly His Thr Leu Gln Val Ala Gly Leu Arg Gly Pro Val Leu
212 640      645      650      655
214 cgc ctg cgg gag ccg ctg ggt gtg ctg gct gtg gtg tgt ccg gac gag 2075
215 Arg Leu Arg Glu Pro Leu Gly Val Leu Ala Val Val Cys Pro Asp Glu
216      660      665      670
218 tgg ccc ctg ctt gcc ttc gtg tcc ctg ctg gct ccc gcc ctg gcc tac 2123
219 Trp Pro Leu Leu Ala Phe Val Ser Leu Leu Ala Pro Ala Leu Ala Tyr
220      675      680      685
222 ggc aac act gtg gtc atg gtg ccc agt gcg gcc tgt cct ctg ctg gcc 2171
223 Gly Asn Thr Val Val Met Val Pro Ser Ala Ala Cys Pro Leu Leu Ala
224      690      695      700
226 ctg gag gtc tgc cag gac atg gcc acc gtg ttc cca gca ggc ctg gcc 2219
227 Leu Glu Val Cys Gln Asp Met Ala Thr Val Phe Pro Ala Gly Leu Ala
228      705      710      715
230 aac gtg gtg aca gga gac cgg gac cat ctg acc cgc tgc ctg gcc ttg 2267
231 Asn Val Val Thr Gly Asp Arg Asp His Leu Thr Arg Cys Leu Ala Leu
232 720      725      730      735
234 cac caa gac gtc cag gcc atg tgg tat ttc gga tca gcc cag ggt tcc 2315
235 His Gln Asp Val Gln Ala Met Trp Tyr Phe Gly Ser Ala Gln Gly Ser
236      740      745      750
238 cag ttt gtc gag tgg gcc tcg gca gga aac ctg aaa ccg gtg tgg gcg 2363
239 Gln Phe Val Glu Trp Ala Ser Ala Gly Asn Leu Lys Pro Val Trp Ala
240      755      760      765
242 agc agg ggc tgc ccg cgg gcc tgg gac cag gag gcc gag ggg gca ggc 2411
243 Ser Arg Gly Cys Pro Arg Ala Trp Asp Gln Glu Ala Glu Gly Ala Gly
244      770      775      780
246 cca gag ctg ggg ctg cga gtg gcg cgg acc aag gcc ctg tgg ctg cct 2459
247 Pro Glu Leu Gly Leu Arg Val Ala Arg Thr Lys Ala Leu Trp Leu Pro
248      785      790      795
250 atg ggg gac tgatgcctga gcgccaccta ctgcattttg gacacctcac 2508
251 Met Gly Asp
252 800
254 accaagggga gatgcacccc acagacacct gggactttcc ccttctggtt cctgtgtctc 2568
255 ccaataaact ctctgaccaa ccctaaaaaa aaaaaaaaaa aaaaaaaaaa rwarmaactt 2628
256 ctggcagata tgaggctttt ttcttttttt tt 2660
258 <210> SEQ ID NO: 2
259 <211> LENGTH: 802

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RAW SEQUENCE LISTING

DATE: 08/10/2005

PATENT APPLICATION: US/10/664,506A

TIME: 13:55:23

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08102005\J664506A.raw

260 <212> TYPE: PRT

261 <213> ORGANISM: Homo sapiens

263 <400> SEQUENCE: 2

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264 Met Ala Ala Thr Arg Ala Gly Pro Arg Ala Arg Glu Ile Phe Thr Ser
265 1 5 10 15
266 Leu Glu Tyr Gly Pro Val Pro Glu Ser His Ala Cys Ala Leu Ala Trp
267 20 25 30
268 Leu Asp Thr Gln Asp Arg Cys Leu Gly His Tyr Val Asn Gly Lys Trp
269 35 40 45
270 Leu Lys Pro Glu His Arg Asn Ser Val Pro Cys Gln Asp Pro Ile Thr
271 50 55 60
272 Gly Glu Asn Leu Ala Ser Cys Leu Gln Ala Gln Ala Glu Asp Val Ala
273 65 70 75 80
274 Ala Ala Val Glu Ala Ala Arg Met Ala Phe Lys Gly Trp Ser Ala His
275 85 90 95
276 Pro Gly Val Val Arg Ala Gln His Leu Thr Arg Leu Ala Glu Val Ile
277 100 105 110
278 Gln Lys His Gln Arg Leu Leu Trp Thr Leu Glu Ser Leu Val Thr Gly
279 115 120 125
280 Arg Ala Val Arg Glu Val Arg Asp Gly Asp Val Gln Leu Ala Gln Gln
281 130 135 140
282 Leu Leu His Tyr His Ala Ile Gln Ala Ser Thr Gln Glu Glu Ala Leu
283 145 150 155 160
284 Ala Gly Trp Glu Pro Met Gly Val Ile Gly Leu Ile Leu Pro Pro Thr
285 165 170 175
286 Phe Ser Phe Leu Glu Met Met Trp Arg Ile Cys Pro Ala Leu Ala Val
287 180 185 190
288 Gly Cys Thr Val Val Ala Leu Val Pro Pro Ala Ser Pro Ala Pro Leu
289 195 200 205
290 Leu Leu Ala Gln Leu Ala Gly Glu Leu Gly Pro Phe Pro Gly Ile Leu
291 210 215 220
292 Asn Val Val Ser Gly Pro Ala Ser Leu Val Pro Ile Leu Ala Ser Gln
293 225 230 235 240
294 Pro Gly Ile Arg Lys Val Ala Phe Cys Gly Ala Pro Glu Glu Gly Arg
295 245 250 255
296 Ala Leu Arg Arg Ser Leu Ala Gly Glu Cys Ala Glu Leu Gly Leu Ala
297 260 265 270
298 Leu Gly Thr Glu Ser Leu Leu Leu Leu Thr Asp Thr Ala Asp Val Asp
299 275 280 285
300 Ser Ala Val Glu Gly Val Val Asp Ala Ala Trp Ser Asp Arg Gly Pro
301 290 295 300
302 Gly Gly Leu Arg Leu Leu Ile Gln Glu Ser Val Trp Asp Glu Ala Met
303 305 310 315 320
304 Arg Arg Leu Gln Glu Arg Met Gly Arg Leu Arg Ser Gly Arg Gly Leu
305 325 330 335
306 Asp Gly Ala Val Asp Met Gly Ala Arg Gly Ala Ala Ala Cys Asp Leu
307 340 345 350
308 Val Gln Arg Phe Val Arg Glu Ala Gln Ser Gln Gly Ala Gln Val Phe
309 355 360 365

```

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/664,506A

DATE: 08/10/2005
TIME: 13:55:24

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF4\08102005\J664506A.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 6,8

Seq#:4; N Pos. 1337

Seq#:7; N Pos. 1606,1620,1631,1655,1658,1666,1673,1688,1705,1711

Seq#:18; Xaa Pos. 96,97,98,99,100,101,102,103,104,105,106,107,108,109,110

Seq#:18; Xaa Pos. 111,112,113,114,162,163,164,165,166,167,168,169,170,171

Seq#:18; Xaa Pos. 172,173,174,175,176,177,178,179,180,181,182,385,386,387

Seq#:18; Xaa Pos. 388,389,390,391,392,393,394,395,396,397,398,399,400,401

Seq#:18; Xaa Pos. 402,403,404,405,406,407,408,409,410,411,412,413,414,415

Seq#:18; Xaa Pos. 416,417,418,419,420,421,422,423,424,425,426,427,428,429

Seq#:18; Xaa Pos. 430,431,432,433,434,435,436,437,438,439,440,441,442,443

Seq#:18; Xaa Pos. 444,445,446,447,448,449,450,488,489,490,491,492,493,494

Seq#:18; Xaa Pos. 495,496,497,498,499,500,516,517,518,519,520,521,522,523

Seq#:18; Xaa Pos. 524,525,526,527,528,529

Seq#:25; Xaa Pos. 139,140,141,142,143,144,145,146,147,148,149,150,192,193

Seq#:25; Xaa Pos. 194,195,196,197,198,199,200,201,202,203,204,205,206,207

Seq#:25; Xaa Pos. 208,209,210,211,212,213,214,215,216,217

VERIFICATION SUMMARY

DATE: 08/10/2005

PATENT APPLICATION: US/10/664,506A

TIME: 13:55:24

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\08102005\J664506A.raw

L:48 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:596 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:1323
L:837 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:1577
M:341 Repeated in SeqNo=7
L:1717 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1724 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1731 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1738 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:80
M:341 Repeated in SeqNo=18
L:2057 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:2064 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:2081 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:128
M:341 Repeated in SeqNo=25

**Raw Sequence Listing before editing,
for reference only**



IFW16

RAW SEQUENCE LISTING

DATE: 08/09/2005

PATENT APPLICATION: US/10/664,506A

TIME: 14:06:52

Input Set : A:\SEQUENCE LISTING.txt

Output Set: N:\CRF4\08092005\J664506A.raw

4 <110> APPLICANT: Meyers, Rachel
 5 Cook, William James
 6 Williamson, Mark
 7 Rudolph-Owen, Laura A.
 8 Gimeno, Ruth
 10 <120> TITLE OF INVENTION: 21481, A NOVEL DEHYDROGENASE MOLECULE
 11 AND USES THEREFOR
 14 <130> FILE REFERENCE: MPI00-079P1RCP2CN1M
 16 <140> CURRENT APPLICATION NUMBER: 10/664,506A
 17 <141> CURRENT FILING DATE: 2003-09-17
 19 <150> PRIOR APPLICATION NUMBER: 09/838,561
 20 <151> PRIOR FILING DATE: 2001-04-18
 22 <150> PRIOR APPLICATION NUMBER: 09/816,760
 23 <151> PRIOR FILING DATE: 2001-03-23
 25 <150> PRIOR APPLICATION NUMBER: 09/634,955
 26 <151> PRIOR FILING DATE: 2000-08-08
 28 <150> PRIOR APPLICATION NUMBER: 60/192,002
 29 <151> PRIOR FILING DATE: 2000-03-24
 31 <160> NUMBER OF SEQ ID NOS: 34
 33 <170> SOFTWARE: FastSEQ for Windows Version 4.0

*Does Not Comply
Corrected Diskette Needed*

ERRORED SEQUENCES

2278 <210> SEQ ID NO: 34
 2279 <211> LENGTH: 52
 2280 <212> TYPE: PRT
 2281 <213> ORGANISM: Artificial Sequence
 2283 <220> FEATURE:
 2284 <223> OTHER INFORMATION: hypothetical protein domain
 2288 <400> SEQUENCE: 34
 2289 Tyr Leu Ala Pro Gly Gln Gly Gln Val Ala Ile Val Thr Gly Gly Ala
 2290 1 5 10 15
 2291 Thr Gly Ile Gly Lys Ala Ile Val Lys Glu Leu Leu Glu Leu Gly Ser
 2292 20 25 30
 2293 Asn Val Val Ile Ala Ser Arg Lys Leu Glu Arg Leu Lys Ser Ala Ala
 2294 35 40 45
 2295 Asp Glu Leu Gln
 2296 50

E--> 2301 36

VERIFICATION SUMMARY

DATE: 08/09/2005

PATENT APPLICATION: US/10/664;506A

TIME: 14:06:54

Input Set : A:\SEQUENCE LISTING.txt

Output Set: N:\CRF4\08092005\J664506A.raw

L:48 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:1
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0
L:596 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:4
L:683 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:1323
L:837 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:7
L:941 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:7 after pos.:1577
M:341 Repeated in SeqNo=7
L:1717 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:1724 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1731 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1738 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:18
L:1749 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:80
M:341 Repeated in SeqNo=18
L:2057 M:281 W: Numeric Fields not Ordered, <221> Sort in ascending order!
L:2064 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:25
L:2081 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25 after pos.:128
M:341 Repeated in SeqNo=25
L:2301 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:34

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- ☐ **SKEWED/SLANTED IMAGES**
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- ☐ **GRAY SCALE DOCUMENTS**
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